SEQUENCE LISTING

<110> YAROVINSKY, TIMUR

<120> TOPOISOMERASE ACTIVATED OLIGONUCLEOTIDE ADAPTORS AND
      USES THEREFOR

<130> UIA-031.01

<140> 09/871,607
<141> 2001-05-31

<150> 60/208,662
<151> 2000-05-31

<160> 12

<170> PatentIn Ver. 2.1

<210> 1
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      oligonucleotide

<400> 1
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gtaccacctc accagtgtct 20

<210> 6
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aaatgatggc cagagacca 19

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acagatgtag tagtatatga acaaacgtgg gaggaggcgt taactagatt aatttttgtg 180
ggaagtgatt caaaaggacg tagacaatac ttttacggaa aaatgcatgt acagaatcgc 240
aacgctaaaa gagatcgtat ttttgttaga gtatataacg ttatgaaacg aattaattgt 300
tttataaaca aaaatataaa gaaatcgtcc acagattcca attatcagtt ggcggttttt 360
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gaaacagtag ggttattaac actaaaaaat aaacacatag aaataagtcc cgatgaaata 480
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tccatatctc ctcttccatc accaaaaaag ttaatagcgt taactatcaa acaaactgct 780
gaagtggtag gtcatactcc atcaatttca aaaagagctt atatggcaac gactatttta 840
gaaatggtaa aggataaaaa ttttttagat gtagtatcta aaactacgtt cgatgaattc 900

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<213> Vaccinia virus

<400> 8

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            20                  25                  30

His Val Lys Ile Pro Thr His Leu Thr Asp Val Val Val Tyr Glu Gln
        35                  40                  45

Thr Trp Glu Glu Ala Leu Thr Arg Leu Ile Phe Val Gly Ser Asp Ser
    50                  55                  60

Lys Gly Arg Arg Gln Tyr Phe Tyr Gly Lys Met His Val Gln Asn Arg
65                  70                  75                  80

Asn Ala Lys Arg Asp Arg Ile Phe Val Arg Val Tyr Asn Val Met Lys
                85                  90                  95

Arg Ile Asn Cys Phe Ile Asn Lys Asn Ile Lys Lys Ser Ser Thr Asp
            100                 105                 110

Ser Asn Tyr Gln Leu Ala Val Phe Met Leu Met Glu Thr Met Phe Phe
        115                 120                 125

Ile Arg Phe Gly Lys Met Lys Tyr Leu Lys Glu Asn Glu Thr Val Gly
    130                 135                 140

Leu Leu Thr Leu Lys Asn Lys His Ile Glu Ile Ser Pro Asp Glu Ile
145                 150                 155                 160

Val Ile Lys Phe Val Gly Lys Asp Lys Val Ser His Glu Phe Val Val
                165                 170                 175

His Lys Ser Asn Arg Leu Tyr Lys Pro Leu Leu Lys Leu Thr Asp Asp
            180                 185                 190

Ser Ser Pro Glu Glu Phe Leu Phe Asn Lys Leu Ser Glu Arg Lys Val
        195                 200                 205

Tyr Glu Cys Ile Lys Gln Phe Gly Ile Arg Ile Lys Asp Leu Arg Thr
    210                 215                 220

Tyr Gly Val Asn Tyr Thr Phe Leu Tyr Asn Phe Trp Thr Asn Val Lys
225                 230                 235                 240

Ser Ile Ser Pro Leu Pro Ser Pro Lys Lys Leu Ile Ala Leu Thr Ile
                245                 250                 255
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Lys Gln Thr Ala Glu Val Val Gly His Thr Pro Ser Ile Ser Lys Arg
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Ala Tyr Met Ala Thr Thr Ile Leu Glu Met Val Lys Asp Lys Asn Phe
275 280 285

Leu Asp Val Val Ser Lys Thr Thr Phe Asp Glu Phe Leu Ser Ile Val
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Val Asp His Val Lys Ser Ser Thr Asp Gly
305 310

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